GenCore version 5.1.6

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OM protein - nucleic search, using frame_plus_p2n model

Run on: December 14, 2003, 14:23:30 ; Search time 86 Seconds
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Title: US-10-021-698-111
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Searched: 569978 seqs, 220691566 residues

Total number of hits satisfying chosen parameters: 1139956

Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
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                 Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

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2	3235	99.3	1853	4	US-09-638-857-19	Sequence 19, Appl
3	2677	82.2	3540	3	US-08-842-079-16	Sequence 16, Appl
4	2677	82.2	3540	4	US-09-638-857-16	Sequence 16, Appl
5	896	27.5	1750	4	US-09-016-434-831	Sequence 831, App
6	896	27.5	1762	2	US-08-742-621-2	Sequence 2, Appli
7	880.5	27.0	1206	3	US-09-191-608-21	Sequence 21, Appl
8	867	26.6	1997	3	US-08-750-134A-6	Sequence 6, Appli
9	867	26.6	1997	3	US-09-363-745-6	Sequence 6, Appli
10	773	23.7	2643	2	US-08-750-134A-10	Sequence 10, Appl
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12	769.5	23.6	577	4	US-09-016-434-829	Sequence 829, App

13	740	22.7	1837	2	US-08-750-134A-4	Sequence 4, Appli
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15	681.5	20.9	1243	3	US-09-191-136-15	Sequence 15, Appl
16	678.5	20.8	1753	2	US-08-750-134A-8	Sequence 8, Appli
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18	669.5	20.5	1421	3	US-09-191-136-13	Sequence 13, Appl
19	656.5	20.2	1421	3	US-09-191-608-14	Sequence 14, Appl
20	656.5	20.2	1436	3	US-09-191-608-13	Sequence 13, Appl
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22	627.5	19.3	1697	3	US-09-381-681-2	Sequence 2, Appli
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24	624.5	19.2	1360	3	US-09-191-136-30	Sequence 30, Appl
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29	328.5	10.1	531	3	US-09-191-608-8	Sequence 8, Appli
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33	103.5	3.2	1704	4	US-09-252-991A-11910	Sequence 11910, A
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37	100.5	3.1	4411529	3	US-09-103-840A-1	Sequence 1, Appli
38	99.5	3.1	2455	4	US-09-227-357-79	Sequence 79, Appl
39	99	3.0	1493	4	US-09-016-434-1175	Sequence 1175, Ap
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c 42	97.5	3.0	2348	4	US-09-393-554-10	Sequence 10, Appl
43	97	3.0	6168	4	US-09-071-035-457	Sequence 457, App
44	97	3.0	6168	4	US-09-071-035-461	Sequence 461, App
45	97	3.0	6168	4	US-09-071-035-465	Sequence 465, App

ALIGNMENTS

RESULT 1
US-08-842-079-19
; Sequence 19, Application US/08842079
; Patent No. 6133434
; GENERAL INFORMATION:
; APPLICANT: BUELL, GARY N.
; APPLICANT: SURPRENANT, ANNMARIE
; APPLICANT: KAWASHIMA, ERIC
; TITLE OF INVENTION: A PURINERGIC RECEPTOR
; FILE REFERENCE: 1430-160
; CURRENT APPLICATION NUMBER: US/08/842,079
; CURRENT FILING DATE: 1997-04-28
; SOFTWARE: PatentIn Ver. 2.0
; NUMBER OF SEQ ID NOS: 20
; SEQ ID NO 19
; LENGTH: 1853
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Synthetic
US-08-842-079-19

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Percent Similarity: 99.33%   Conservative: 0
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Query Match: 99.29%   Indels: 0
DB: 3   Gaps: 0

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; Sequence 19, Application US/09638857
; Patent No. 6509163
; GENERAL INFORMATION:
; APPLICANT: BUELL, GARY N.
; APPLICANT: SURPRENANT, ANNMARIE
; APPLICANT: KAWASHIMA, ERIC
; TITLE OF INVENTION: A PURINERGIC RECEPTOR
; FILE REFERENCE: 1430-160
; CURRENT APPLICATION NUMBER: US/09/638,857
; CURRENT FILING DATE: 2000-08-15
; PRIOR APPLICATION NUMBER: 08/842,079
; PRIOR FILING DATE: 1997-04-28
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; OTHER INFORMATION: Description of Artificial Sequence:Synthetic
US-09-638-857-19

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Qy 61   ValHisThrLysValLysGlyIleAlaLeuValLysGluLeuValAlaAsnGlyVal 80
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OM nucleic - nucleic search, using sw model

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                  GenCore version 5.1.6
         Copyright (c) 1993 - 2003 Compugen Ltd.
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Searched: 56978 seqs, 22069156 residues

Total number of hits satisfying chosen parameters: 1139956

Minimum DB seq length: 0
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and is derived by analysis of the total score distribution.

SUMMARIES

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| 2 | 1834.6 | 36.1 | 1853 | 4 | US-09-638-857-19 | Sequence 19, Appl |
| 3 | 1285.8 | 25.3 | 3540 | 3 | US-08-842-079-16 | Sequence 16, Appl |
| 4 | 1285.8 | 25.3 | 3540 | 4 | US-09-638-857-16 | Sequence 16, Appl |
| 5 | 450 | 8.8 | 577 | 4 | US-09-016-434-829 | Sequence 829, App |
| 6 | 383.2 | 7.5 | 59065 | 4 | US-09-813-817-3 | Sequence 3, Appli |
| 7 | 383.2 | 7.5 | 59065 | 4 | US-09-978-197-3 | Sequence 3, Appli |
| c 8 | 332 | 6.5 | 55298 | 4 | US-09-491-356C-1 | Sequence 1, Appli |
| c 9 | 330.8 | 6.5 | 36651 | 4 | US-09-738-894A-3 | Sequence 3, Appli |
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| 11 | 328.4 | 6.5 | 15977 | 3 | US-09-608-285A-59 | Sequence 59, Appl |
| c 12 | 326.8 | 6.4 | 14747 | 4 | US-09-608-285A-42 | Sequence 42, Appl |
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| 16 | 311.2 | 6.1 | 87350 | 4 | US-09-618-166-79 | Sequence 79, Appl |
| 17 | 311.2 | 6.1 | 87543 | 4 | US-09-791-211-3 | Sequence 3, Appli |
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| 24 | 306 | 6.0 | 31571 | 1 | US-08-323-443B-1 | Sequence 1, Appli |
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| c 41 | 293.8 | 5.8 | 6769 | 5 | PCT-US95-10202-20 | Sequence 20, Appl |
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| c 43 | 293.8 | 5.8 | 6769 | 5 | PCT-US95-10220-20 | Sequence 20, Appl |
| c 44 | 293.2 | 5.8 | 7676 | 1 | US-08-451-777A-7 | Sequence 7, Appli |
| c 45 | 293.2 | 5.8 | 7676 | 2 | US-08-451-778A-7 | Sequence 7, Appli |

ALIGNMENTS

RESULT 1
US-08-842-079-19

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; Sequence 19, Application US/08842079
; Patent No. 6133434
; GENERAL INFORMATION:
; APPLICANT: BUELL, GARY N.
; APPLICANT: SURPRENANT, ANNMARIE
; APPLICANT: KAWASHIMA, ERIC
; TITLE OF INVENTION: A PURINERGIC RECEPTOR
; FILE REFERENCE: 1430-160
; CURRENT APPLICATION NUMBER: US/08/842,079
; CURRENT FILING DATE: 1997-04-28
; NUMBER OF SEQ ID NOS: 20
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 19
; LENGTH: 1853
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Synthetic
US-08-842-079-19
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Qy   408  CCAAGAGCAGCGGTTGTGTCCCGAGTATCCACCCGGACGAGGACGCTCGTTCCTCTGACCG  467
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   365  CCAAGAGCAGCGGTTGTGTCCCGAGTATCCACCCGGACGAGGACGCTCGTTCCTCTGACCG  424

Qy   468  AGGTTGTAAAAAGGATGGATGGACCCGCAGAGCAAAGGAATTCAGACCGGAAGGTGTGT   527
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   425  AGGTTGTAAAAAGGATGGATGGACCCGCAGAGCAAAGGAATTCAGACCGGAAGGTGTGT   484

Qy   528  AGTGCATGAAGGGAACCAGAAGACTTGTGAAGTCTCTGCCTGGTGCCCATCGAGGCAGT   587
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   485  AGTGCATGAAGGGAACCAGAAGACTTGTGAAGTCTCTGCCTGGTGCCCATCGAGGCAGT   544

Qy   588  GGAAGAGGCCCCCCGGCCTGCTCTTGACAGTGCCGAAAACTTCACTGTGCTCATCAA   647
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   545  GGAAGAGGCCCCCCGGCCTGCTCTTGACAGTGCCGAAAACTTCACTGTGCTCATCAA   604

Qy   648  GAACAATATGCACTTCCCCGGCCACAACTACACCACGAGAAACATCCTGCCAGTTTAAA   707
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   605  GAACAATATCGACTTCCCCGGCCACAACTACACCACGAGAAACATCCTGCCAGTTTAAA   664

Qy   708  CATCACTTGTACCTTCCACAAGACTCAGAATCCACAGTGTCCCATTTTCCGACTAGGAGA   767
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   665  CATCACTTGTACCTTCCACAAGACTCAGAATCCACAGTGTCCCATTTTCCGACTAGGAGA   724

Qy   768  CATCTTCCGAGAAACAGGCGATAATTTTCAGATGGCAATTCAGGGGCGGAATAATGGG   827
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   725  CATCTTCCGAGAAACAGGCGATAATTTTCAGATGGCAATTCAGGGGCGGAATAATGGG   784

Qy   828  CATTGAGAATCTACTGGGACTGCAACCTAGACCGTTGGTTCCATCACTGCCGTCCAAATA   887
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   785  CATTGAGAATCTACTGGGACTGCAACCTAGACCGTTGGTTCCATCACTGCCGTCCAAATA   844

Qy   888  CAGTTTCCGTCGCCTTGACGACAAGACCACCAACGTGTCCTTGTACCCTGGCTACAACTT   947
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   845  CAGTTTCCGTCGCCTTGACGACAAGACCACCAACGTGTCCTTGTACCCTGGCTACAACTT   904

Qy   948  CAGATACGCCAAGTACTACAAGGAAAACAATGTTGAGAAACGGACTCTGATAAAAGTCTT   1007
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   905  CAGATACGCCAAGTACTACAAGGAAAACAATGTTGAGAAACGGACTCTGATAAAAGTCTT   964

Qy   1008  CGGGATCCGTTTTGACATCCTGGTTTTTGGCACCGGAGGAAAATTTGACATTATCCAGCT  1067
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   965  CGGGATCCGTTTTGACATCCTGGTTTTTGGCACCGGAGGAAAATTTGACATTATCCAGCT  1024

Qy   1068  GGTTGTTGTACATCGGCTCAACCCTCTGGTCTGGCCACTGTTGTTCATCGACTT  1127
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1025  GGTTGTTGTACATCGGCTCAACCCTCTCCTACTTGGTCTGGCCGCTGTTCATCGACTT  1084

Qy   1128  CCTCATCGACACTTACTCCAGTAACTGCTGTCGCTGTCGCTCCCATATTATCCCTGGTGTCAAGTG  1187
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1085  CCTCATCGACACTTACTCCAGTAACTGCTGTCGCTGTCGCTCCCATATTTATCCCTGGTGTCAAGTG  1144

Qy   1188  CTGTTGTGTACATCGGTGGTCAACGAATACTACTACCAGGAGAAGTGCGAGTCCATTGTGGA  1247
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1145  CTGTTGTGTACATCGGTGGTCAACGAATACTACTACAGGAGAAGTGCGAGTCCATTGTGGA  1204

Qy   1248  GCCAAAGCCGACATTAAAGTATGTGTCCTTTGTGGATGAATCCCACATTAGGATGGTGAA  1307
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1205  GCCAAAGCCGACATTAAAGTATGTGTCCTTTGTGGATGAATCCCACATTAGGATGGTGAA  1264

Qy   1308  CCAGCAGCTACTAGGGAGAAGTTTGCAAGATGTCAAGGGCCAAGAAGTCCCAAGACCTGC  1367
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1265  CCAGCAGCTACTAGGGAGAAGTTTGCAAGATGTCAAGGGCCAAGAAGTCCCAAGACCTGC  1324

Qy   1368  GATGGACTTCACAGATTTGTCCCAGGCTGCCCTCCATGCACCCCGATTCC  1427
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1325  GATGGACTTCACAGATTTGTCCCAGGCTGCCCTCCATGCACACACCCCGATTCC  1384

Qy   1428  TGGACAACCAGGAGGAGATCAGCTGCTTAGAAAGGAGGCGACTTCCTAGATCCAGGGATAG  1487
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1385  TGGACAACCAGAGGAGATCAGCTGCTTAGAAAGGAGGCGACTCCTAGATCCAGGGATAG  1444

Qy   1488  CCCGTCTGGTGCCCAGTGTGGAAGATGCCTCCCATCTCAACTCCCTGAGCCACAGGTG  1547
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Db   1445  CCCCGTCTGTGCCCAGTGTGGCAGTGGGAGCTGCCTCCCATCTCAACTCCCTGAGCCACCTTCAGAGCTGTT  1504
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Qy   1548  CCTGGAGGAGCTGGTCTGTCCGGAAAAAGCCGGGGGCTCCTCCGCATCACCTCACCACCTCAGAGCTGTT  1607
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1505  CCTGGAGGAGCTGGTCTGTCCGGAAAAAGCCGGGGGCTCCTCCGCATCACCTCACCACCTCAGAGCTGTT  1564

Qy   1608  CAGGAAGCTGGTCCTGTCCAGACACGTCCTCCTCGCTCCTGCTCTTCTACCAGGAGCCCT  1667
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1565  CAGGAAGCTGGTCCTGTCCAGACACGTCCTCCTCGCTCCTGCTCTTCTACCAGGAGCCCT  1624

Qy   1668  GCTGGCGCTGGATGTGATTCACCAACAGCCGGCTGCGGCACTGTGCGCTACAGGTGCTA  1727
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1625  GCTGGCGCTGGATGTGATTCACCAACAGCCGGCTGCGGCACTGTGCGCTACAGGTGCTA  1684

Qy   1728  CGCCACCTGGCCGCTTCGGCTCGGCTTCGCTCCCAGGACCATCAGGTGCCATCCTGCCCCAGCTGCTG  1787
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1685  CGCCACCTGGCCGCTTCGGCTCGGCTTCGCTCCCAGGACCATCAGGTGCTGACTTTGCCATCCTGCCCCAGCTGCTG  1744

Qy   1788  CCGCTGAGGGATCCGGAAAGAGTTTCAAAGAGTGAAGGGCAGTACAGTGGCTTCAAGAG  1847
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1745  CCGCTGAGGGATCCGGAAAGAGTTTCCGAAGAGTGTAGGGCAGTAGGGCAGTGGCTTCAAGAG  1804

Qy   1848  TCCTTACTGAAGCCAGGACACCGTGGCTCACGTCGTAATCCGCGCTT  1896
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1805  TCCTTACTGAAGCCAGGCAGCACCGTGGCTCACGTCTGTAATCCCCACCTTTT  1853
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RESULT 2
US-09-638-857-19
; Sequence 19, Application US/09638857
; Patent No. 6509163
; GENERAL INFORMATION:
; APPLICANT: BUELL, GARY N.
; APPLICANT: SURPRENANT, ANNMARIE
; APPLICANT: KAWASHIMA, ERIC
; TITLE OF INVENTION: A PURINERGIC RECEPTOR
; FILE REFERENCE: 1430-160
; CURRENT APPLICATION NUMBER: US/09/638,857
; CURRENT FILING DATE: 2000-08-15
; PRIOR APPLICATION NUMBER: 08/842,079
; PRIOR FILING DATE: 1997-04-28
; NUMBER OF SEQ ID NOS: 20
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 19
; LENGTH: 1853
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Synthetic
US-09-638-857-19

Query Match   36.1%;   Score 1834.6;   DB 4;   Length 1853;
Best Local Similarity 99.5%; Pred. No. 0;
Matches 1840; Conservative 0; Mismatches 9; Indels 0; Gaps 0;

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Qy   48   CGCAGGAGGAGGAGGCTGTCACCATGCCGCCTGCTCGCAGCTGCAGTGATGTGATGTTTCCAGTA  107
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5    CGCAGGAGGAGGAGGCTGTCACCATGCCGCCTGCTCGCAGCTGCAGTGATGTGATGTTTCCAGTA  64

Qy   108  TGAGACGAACAAAGTCACTCCGATCCAGAGCAAGATGAATATGGCACCATTAAGTGGTTCTT  167
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   65   TGAGACGAACAAAGTCACTCCGATCCAGAGCAAGATGAATATGGCACCATTAAGTGGTTCTT  124

Qy   168  CCACGTGATCATCTTTTCCTACGTTTGCTTGGTGGTGGTGAGTGCAAGCTGTAACCAGCG  227
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   125  CCACGTGATCATCTTTTCCTACGTTTGCTTGGTGGTGGTGAGTGCAAGCTGTAACCAGCG  184

Qy   228  GAAAGAGCCTGTCACAGTTCTGTGCACACCAAGGTGAAGGGATGCAGAGGTGAAAGA  287
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   185  GAAAGAGCCTGTCACAGTTCTGTGCACACCAAGGTGAAGGGATGCAGAGGTGAAAGA  244

Qy   288  GGAGATCGTGAGAATGGAGTGAAGAAGTTGGTGCACACAGTGTCTTTGACACCGCAGACTA  347
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